



## *Anopheles (Cellia) rampae* n. sp., alias chromosomal form K of the Oriental Maculatus Group (Diptera: Culicidae) in Southeast Asia

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### Abstract

Chromosomal form K (also known as species K) of the *Anopheles maculatus* group of sibling species in the Oriental Region is diagnosed and formally named *An. (Cellia) rampae* Harbach & Somboon, n. sp. The male genitalia, pupa and fourth-instar larva are illustrated and information is provided on the morphology, systematics, bionomics and distribution of the species.

**Key words:** *Anopheles rampae*, *Anopheles maculatus*, mosquitoes, Neocellia Series, new species

### Introduction

The Maculatus Group is an assemblage of eight formally named Oriental mosquito species within the Neocellia Series of *Anopheles* subgenus *Cellia* in the Old World. *Anopheles maculatus* Theobald was first recognised as a complex of species based on polytene chromosome rearrangements (Green & Baimai, 1984; Green *et al.*, 1985), and later confirmed by metaphase karyotypes (Baimai *et al.*, 1993) and DNA sequence data (Ma *et al.*, 2002, 2006; Walton *et al.*, 2007; Morgan *et al.*, 2009). Correlation of morphological differences with the cytogenetic forms enabled Rattanaarithikul & Green (1987) and Rattanaarithikul & Harbach (1991) to assign both existing (available) and new Latin names to the eight forms, i.e. *An. dispar* Rattanaarithikul & Harbach, *An. dravidicus* Christophers, *An. greeni* Rattanaarithikul & Harbach, *An. maculatus* Theobald, *An. notanandai* Rattanaarithikul & Green, *An. pseudowillmori* (Theobald), *An. sawadwongporni* Rattanaarithikul & Green and *An. willmori* (James). A ninth cytogenetic form, first referred to as species K by Baimai (1989), has only recently been embraced as a species of the Group based on an accumulation of evidence that includes morphology (Harbach, 2004; Rattanaarithikul *et al.*, 2006), mitotic karyotype (Baimai, 1989; Baimai *et al.*, 1993), rDNA and mtDNA sequences (Ma *et al.*, 2006; Walton *et al.*, 2007; Morgan *et al.*, 2009) and crossmating studies (Somboon *et al.*, 2008; Thongwat, 2008; Thongwat *et al.*, 2008). Based on this body of evidence, species K is described and named *An. rampae* n. sp. in this report.

### Material and methods

This study is based on specimens with associated larval and pupal exuviae that were reared from cytotyped progeny broods of females and fourth-instar larvae collected at various localities in Chaiyaphum, Loei, Mukdahan, Nakhon Phanom, Phetchabun, Nong Khai, Ubon Ratchathani and Udon Thani Provinces of northeastern Thailand. Observations of adults were made under artificial light. Larval and pupal chaetotaxy were studied using bright-field microscopy. Setal branching counts were obtained from 10 specimens (20 setae) of each life stage. Unless indicated otherwise, numbers in parentheses represent modes of the reported ranges. The morphological terminology

and abbreviations contained in the Anatomical Glossary of the online Mosquito Taxonomic Inventory (<http://mosquito-taxonomic-inventory.info/anatomical-glossary-overview>) are used in the descriptions and illustrations, respectively. The symbols ♀, ♂, Le, Pe and L used in the **Type Series** represent adult female(s), adult male(s), larval exuviae, pupal exuviae and fourth-instar larvae, respectively.

## Taxonomy

### *Anopheles (Cellia) rampae* Harbach & Somboon, n. sp.

*Anopheles maculatus* form K of Baimai *et al.*, 1993: 116, 117, 119–121 (Thailand; metaphase karyotype); Harbach, 2004: 548 (notes); Rattarithikul *et al.*, 2005: 29 (Thailand; species list); Ma *et al.*, 2006: 274, 275, 277, 278 (ITS2 rDNA, phylogenetic relationships); Rattarithikul *et al.*, 2006: 27, 32, 45, 77, 125 (Thailand; ♀ L keys); Walton *et al.*, 2007: 93, 94, 96–100 (Cambodia, Thailand, Vietnam; ITS2 rDNA); Manguin *et al.*, 2008: 496, 497 (Thailand; review); Somboon *et al.*, 2008: 1317–1321 (Thailand; crossmating); Thongwat, 2008 (Thailand; genetics, morphology); Thongwat *et al.*, 2008: 194–201 (Thailand; crossmating).

*Anopheles maculatus* species K of Baimai, 1989: 147, 152, 157, 159 (Thailand); World Health Organization, 2007: 5, 49, 53–55 (Thailand; review); Morgan *et al.*, 2009: 591, 292, 294–296, 298 (Cambodia, Thailand, Vietnam; mtDNA, rDNA, phylogenetic relationships).

**Diagnosis.** Members of the Maculatus Group are medium-sized mosquitoes that lack upper proepisternal setae, have speckled legs and hindtarsomere 5 entirely pale-scaled. Based on the studies of Rattarithikul & Green (1987) and Rattarithikul *et al.* (2006), the presence of pale scales on most or all of abdominal terga II–VII and vein R2 shorter than vein R2+3 distinguish the adult females of *An. rampae* from those of *An. pseudowillmori*. Despite overall similarity, clusters of dark scales on the posterolateral corners of abdominal terga II–VII and the presence of two dark spots on vein R2+3 of at least one wing (or the presector dark spot the same length on vein R as on the costa and subcosta if only one spot is present) readily distinguish *An. rampae* females from *An. dravidicus*, *An. maculatus* and *An. willmori*. Females of *An. rampae* are usually distinguished from those of *An. notanandai* by having the median pale spot on veins M1, M2 and M3+4 more than twice as long as the dark spot on either side and the furcation of vein R2+3 within the proximal half of the preapical dark spot on vein R1. Females of *An. rampae* and *An. sawadwongporni* are indistinguishable but their eggs are usually distinguished by the greater length and incomplete frill in the former species. Larvae of the Maculatus Group normally have seta 6-III with 20 or more branches. The length of the basal stem of seta 4-M is the only character that distinguishes larvae of *An. rampae* from those of *An. dravidicus*, *An. maculatus*, *An. pseudowillmori* and *An. willmori*. The stem of this seta is no longer than four times its width in *An. rampae* and five or more times its width in the latter four species. The larvae of *An. rampae*, *An. notanandai* and *An. sawadwongporni* are indistinguishable.

The unique metaphase karyotype of *An. rampae* readily distinguishes this species from the other members of the Maculatus Group: chromosome X<sub>2</sub> is metacentric, one arm is entirely heterochromatic whereas the opposite arm consists of a euchromatic portion and centromeric heterochromatin of approximately equal length; chromosome X<sub>3</sub> is submetacentric, the short arm consists of a euchromatic portion and a large block of centromeric heterochromatin similar to that of the X<sub>2</sub> and the heterochromatic long arm is larger than that of the X<sub>2</sub>; chromosome Y<sub>1</sub> is acrocentric or a small submetacentric and Y<sub>2</sub> is very large and submetacentric (Baimai *et al.*, 1993).

Sequence data for the ITS2 region of ribosomal DNA and the COII and ND5 genes of mitochondrial DNA distinguish *An. rampae* from the other species of the Maculatus Group (except *An. notanandai* Rattarithikul & Green which has not been sequenced) (Ma *et al.*, 2006; Walton *et al.*, 2007; Morgan *et al.*, 2009; as form K).

**Female.** *Head:* Proboscis length 1.78–1.88 mm, 1.10–1.29 length of forefemur; maxillary palpus length 1.63–1.83 mm, 0.88–0.97 length of proboscis; palpomeres 2 and 3 black-scaled except at apices, palpomere 3 infrequently with dorsomesal patch of pale scales, preapical black band 0.40–0.77 length of subapical white band and 0.32–0.56 length of apical white band; subapical white band 0.57–1.00 length of apical white band. *Thorax:* Integument dark brown; central area of scutum with narrow white spatulate scales, scales on scutal fossa shorter and broader; scutellum with similar scales. *Wing* (Table 1): Preapical dark spot on costa 1.20–2.30 (mean 1.76) length of preapical pale spot; presector dark spot on vein R 0.71–1.15 (mean 0.91) length of corresponding dark spot on costa; vein R2 short, 1.09–1.77 length of vein R2+3; R2+3 with 2 dark spots on at least one wing (85.5%), distal spot near furcation rarely absent, the 2 spots occasionally joined to form a single large spot; vein R4+5 occasionally

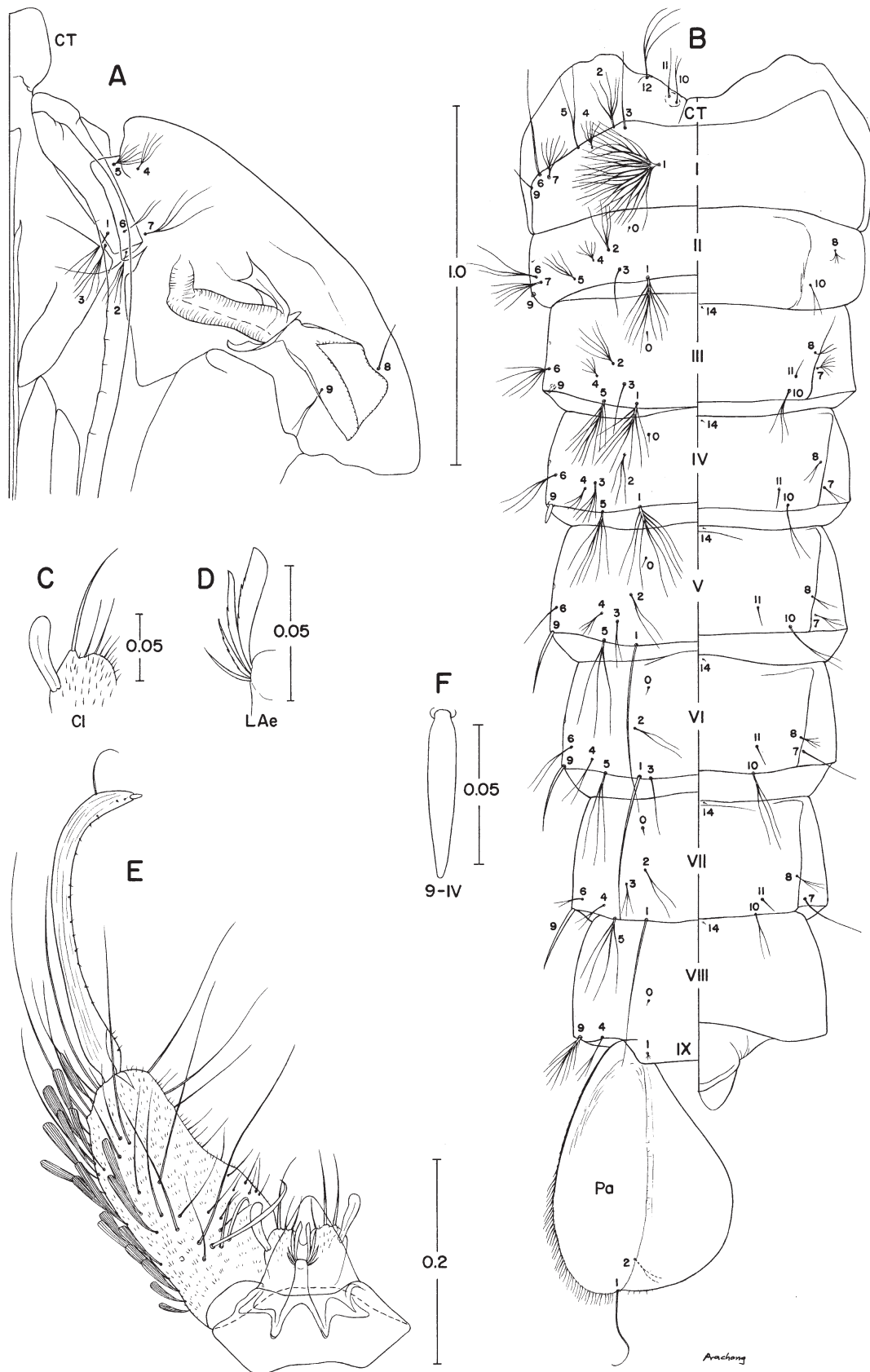
with median dark spot (on one wing in 8.1% of specimens; on 2 wings in 3.2% of specimens); vein M1 with median pale spot, 0.75–2.3 (mean 1.37) length of dark spot on either side; vein M2 with median pale spot, 0.62–2.60 (mean 1.46) length of dark spot on either side; vein M3+4 with long median pale spot, 1.25–2.67 (mean 1.83) length of dark spot on either side; vein 1A with short to long median dark spot, 0.23–0.80 (mean 0.49) length of pale spot on either side, dark fringe spots usually present before apex of vein R1 and between apices of veins R2 and R3. *Legs*: Anterior surface of forecoxa with patch of pale and dark scales at base; femora and tibiae with scattered pale spots and narrow pale patch and/or fringe at apex, ventral surface of forefemur with indefinite stripe of pale scales apically, mid- and hindfemora with indefinite stripe of pale scales ventrally at base; tarsi with pale bands and spots, foretarsomere 1 with 3–7 posterodorsal pale spots, foretarsomere 2 sometimes with a single pale spot in median dark band; midtarsomere 1 with 3–7 posterodorsal pale spots, midtarsomeres 1–3 with narrow pale spot dorsally at apex, midtarsomeres 2 and 3 without median pale spots, midtarsomeres 3 and 4 occasionally with posterodorsal pale spot at base; hindtarsomere 1 with 6–10 posterodorsal pale spots, hindtarsomere 2 sometimes with pale spot in median dark band. *Abdomen*: Integument dark brown, covered with dark setae; terga II–IV sparsely or densely covered with pale spatulate scales posteriorly, posterolateral corners of terga III and IV usually with few black spatulate scales; terga V–VIII largely covered with pale spatulate scales, also with conspicuous patches of black spatulate scales on posterolateral corners; sterna IV–VII with few scattered pale spatulate scales (occasionally on sternum III as well) and median patch of black spatulate scales on posterior margins; sternum VIII with pale falcate and/or spatulate scales laterally.

**TABLE 1.** Wing characteristics of *Anopheles rampae*.

Characteristics	Females (n = 62)		Males (n = 61)	
	One wing (%)	Both wings (%)	One wing (%)	Both wings (%)
Presector dark spot on vein R as long as presector dark spots on subcosta and costa	54.8	40.3	70.5	54.1
Vein R2+3 with two dark spots	85.5	64.5	86.9	73.8
Vein R4+5 with 3 dark spots	8.1	3.2	11.5	4.9
Vein M1 with median pale spot >2 times length of dark spot on either side	19.4	8.1	50.8	23.0
Vein M2 with median pale spot >2 times length of dark spot on either side	72.6	53.2	83.6	63.9
Vein M3+4 with median pale spot >2 times length of dark spot on either side	72.6	50.0	68.9	52.5
Furcation of vein R2+3 within proximal 0.5 of preapical dark spot on vein R1	48.4	29.0	0	0

**Male.** Like female except as follows. *Head*: Proboscis length 2.08–2.18 mm, 1.31–1.50 length of forefemur; maxillary palpus length 2.08–2.03 mm, 0.95–1.02 length of proboscis. *Wing*: Characteristics compared with those of female in Table 1. *Legs*: Foretarsomeres 2 and 3 with apical pale patch or band only. *Abdomen*: Posterolateral corners of terga III and VII occasionally with few black spatulate scales; tergum VIII (ventral in position) with variable amount of pale scaling, with median patch of black spatulate scales posteriorly; sternum VIII (dorsal in position) largely pale-scaled. *Genitalia* (Fig. 1C–E): Gonocoxite with scales on lateral surface, with 4 or 5 slender and lanceolate parabasal setae; gonostylus longer than gonocoxite, with prominent subapical tergo-lateral setae and row of minute setae along sternolateral margin; gonostylar claw short, pigmented. Claspette with tergo-lateral club formed of fused stems and 1 or 2 slender apical setae longer than club. Aedeagus narrow, apex usually with 4 flattened leaflets on either side, distal leaflets serrate on outer margin. Proctiger membranous, lightly sclerotised laterally.

**Egg.** Length 0.49–0.54 mm, width 0.12–0.15 mm; deck usually complete; frill incomplete in middle on both sides, distance between anterior and posterior frills 0.20–0.33 length of deck; ratio of width to length of part of deck enclosed by anterior frill 0.14–0.27, ratio of width between floats to length of part of deck between anterior and posterior frills 0.23–0.62, ratio of width to length of part of deck enclosed by posterior frill 0.16–0.31; float with 14–20 ridges, confluent with median part of deck (part without frill).



**FIGURE 1.** Pupa and male genitalia of *Anopheles (Cellia) rampae*. A, Pupa, left side of cephalothorax, dorsal to right. B, Pupa, dorsal (left) and ventral (right) aspects of metathorax and abdomen. C–E, Male genitalia (tergal aspects): C, apex of claspette; D, apex of aedeagus; E, phallosome and left gonocoxite. Cl, claspette; CT, cephalothorax; LAe, leaflets of aedeagus; Pa, paddle; I–IX = abdominal segments I–IX; 1–14 = setal numbers for specified areas, e.g. seta 3-I. Scales in mm.

**Pupa** (Fig. 1A,B). Character and positions of setae as figured; range and modal number of branches in Table 2. *Cephalothorax*: Trumpet length 0.35–0.43 mm, width 0.08–0.11 mm, index 3.64–4.00. *Abdomen*: Seta 9-II,III very short, blunt; 9-IV–VII long, sharply pointed; ratio of length of seta 9-III/9-IV 0.30–0.55 (mean 0.37), 9-IV/9-V 0.26–0.38 (mean 0.30); 9-VIII with 9–19(13) branches. *Paddle*: Seta 1-P generally shorter, 0.31–0.44 length of paddle.

**TABLE 2.** Range (mode) of numbers of branches for pupal setae of *Anopheles rampae*.

Seta no.	Cephalothorax CT	Abdominal segments									Paddle Pa
		I	II	III	IV	V	VI	VII	VIII	IX	
0	–	–	1,2(1)	1	1	1	1,2(1)	1	1	–	–
1	2–4(3)	nc	4–9(5)	4–10(6)	2–8(3)	1,2(1)	1,2(1)	1,2(1)	–	3,4(4)	1
2	2–4(3)	3–8(4)	3–7(4)	3–7(5)	2–4(3)	2–4(3)	2,3(2)	1–3(2)	–	–	1–3(2)
3	2–4(3)	1,2(1)	1–3(1)	1–3(2)	1–6(4)	1–3(2)	1–3(1)	1–3(2)	–	–	–
4	2–6(3)	2–6(3)	1–7(4)	1–6(3)	1–4(3)	1–5(3)	1–3(1)	1–3(1)	1–3(2)	–	–
5	4–8(5)	1–4(3)	2–6(4)	5–11(7)	3–10(5)	2–7(4)	3–6(4)	3–6(4)	–	–	–
6	2–4(3)	1–5(3)	3–6(3)	2–8(3)	1–4(3)	1–4(1)	1–3(1)	1–3(1)	–	–	–
7	3,4(3)	2–7(3)	3–6(3)	2–6(4)	1–6(3)	1–5(2)	1–3(1)	1,2(1)	–	–	–
8	1–3(1)	–	3–5(4)	1–5(3)	1–3(1)	1,2(1)	1–4(2)	1–4(3)	–	–	–
9	2–4(3)	1,2(1)	1	1	1	1	1	1	9–19(13)	–	–
10	1,2(2)	–	1,2(1)	1–4(2)	1,2(1)	1–3(2)	1–3(2)	1–4(2)	–	–	–
11	1–5(3)	–	–	1,2(1)	1	1,2(1)	1	1,2(1)	–	–	–
12	1–5(3)	–	–	–	–	–	–	–	–	–	–
14	–	–	–	1	1	1	1	1	1	–	–

nc = not counted.

**Larva, fourth-instar** (Fig. 2). Character and placement of setae as figured; range and modal number of branches in Table 3. *Head*: Seta 4-A with 2,3(2) branches; seta 2-C single, with 5–12 short lateral aciculae. *Thorax*: Seta 4-M with 3–5(4) branches, width of basal stem 0.20–0.40 (mean 0.28) of its length; seta 3-T with 1–4(4) branches. *Abdomen*: Seta 1-I not palmate, with 4–7(5) branches; seta II–VII palmate; seta 1-II with 8–12(10) weakly developed leaflets, 1-III with 13–18(16) fully developed leaflets, filaments usually <0.25 length of blades; leaflets of seta 1-IV–V each with relatively long filament, 0.19–0.38 length of blade; seta 4-I with 4–10(7) branches; seta 5-VII with 7–13(9) branches. Pecten plate with 4–6 long and 6–10 short spines.

**DNA sequence.** Ma *et al.* (2006) and Walton *et al.* (2007) reported that the nucleotide sequence of the ITS2 region of ribosomal DNA in *An. rampae* (as form K) is unique among members of the Maculatus Group (*An. notanandai* not included), differing by 3.7% (Walton *et al.*, 2007) from *An. sawadwongporni* Rattanakrithikul & Green, the most closely related species. More recently, Morgan *et al.* (2009) also found a strong sister relationship between *An. rampae* (as form K) and *An. sawadwongporni* based on a Bayesian analysis of the mitochondrial COII and ND5 genes.

**Etymology.** This species is named in honor of Dr Rampa Rattanakrithikul (formerly of the Armed Forces Research Institution of Medical Sciences, Bangkok) for her many important contributions to our knowledge of mosquitoes in Southeast Asia, especially her studies of the Maculatus Group (Rattanakrithikul & Green, 1987; Rattanakrithikul & Harbach, 1991) that provided the taxonomic foundation for further studies of this medically important group of insects.

**Systematics.** The correlation of morphological differences with paracentric inversions of polytene chromosomes and heterochromatin variation in mitotic chromosomes allowed Rattanakrithikul & Green (1987) and Rattanakrithikul & Harbach (1991) to give formal Latin names to eight members of the Maculatus Group. Polytene chromosomes have not been studied in the ninth member of the Group, i.e. *An. rampae*, but as noted above the unique metaphase karyotype readily distinguishes this form from the other species. In particular, the X and Y chromosomes of *An. rampae* are different from those of *An. notanandai* and *An. sawadwongporni*, the most morphologically similar species.



Mitotic karyotypes from brain ganglia of fourth-instar larvae of *An. rampae* (form K) and *An. maculatus* forms B and E are generally similar, consisting of three types of X chromosomes (small submetacentric X<sub>1</sub>, metacentric X<sub>2</sub> and large submetacentric X<sub>3</sub>) and two types of Y chromosomes (acrocentric or small submetacentric Y<sub>1</sub> and large submetacentric Y<sub>2</sub>) (Green *et al.*, 1985; Baimai *et al.*, 1993). Previously, Baimai *et al.* (1993) reported that X<sub>3</sub>, Y<sub>1</sub> and Y<sub>2</sub> occur in form B; X<sub>1</sub>, X<sub>3</sub> and Y<sub>1</sub> occur in form E; and X<sub>2</sub>, X<sub>3</sub> and Y<sub>2</sub> occur in form K. However, Thongwat (2008) found that X<sub>2</sub> and Y<sub>2</sub> occur in all three forms.

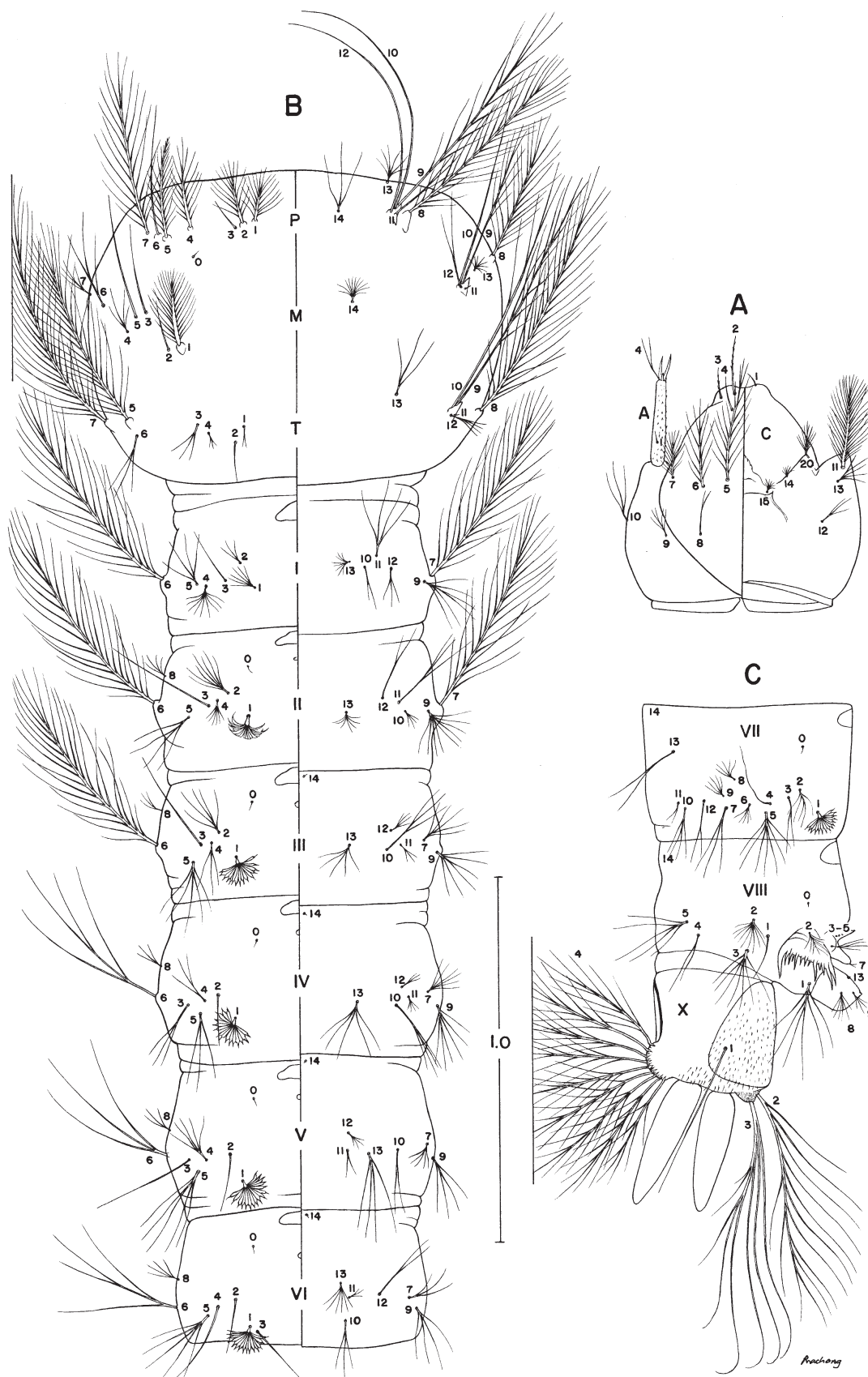
**TABLE 3.** Range (mode) of numbers of branches for fourth-instar larval setae of *Anopheles rampae*.

Seta no.	Head	Thorax			Abdominal segments	
	C	P	M	T	I	II
0	1	1	–	–	–	1
1	1	15–23(19)	26–46(40)	1–3(3)	4–7(5)	8–12(10)
2	1	14–22(19)	1–3(2)	1	3–6(3)	5–7(6)
3	1	1	1	1–4(4)	1,2(1)	1
4	1	13–19(16)	3–5(4)	2–5(4)	4–10(7)	5–11(7)
5	11–22(15)	27–49(40)	1	23–40(30)	4–6(5)	5–7(6)
6	11–21(19)	1	3–6(4)	2–5(3)	24–32(30)	21–32(29)
7	17–25(20)	16–24(22)	2–4(3)	21–36(33)	24–31(27)	24–36(30)
8	1,2(1)	27–39(35)	14–22(20)	29–42(35)	–	2–4(3)
9	1–6(4)	11–16(14)	8–15(11)	12–22(19)	5–7(5)	7–9(8)
10	1–4(3)	1	1	11–19(13)*	1–3(2)	3–5(3)
11	34–41(39)	2–5(3)	1,2(1)	1	3–5(4)	1–3(2)
12	3–6(3)	1	1,2(2)	2–6(4)	3–5(4)	2–4(3)
13	4–7(6)	2–7(5)	5–11(6)	3,4(3)	4–8(7)	5–8(7)
14	nc	3–5(4)	7–11(10)	–	–	–
15	nc	–	–	–	–	–

continued.

Seta no.	Abdominal segments						
	III	IV	V	VI	VII	VIII	X
0	1	1	1	1	1	1	–
1	13–18(16)	12–18(15)	13–17(14)	11–16(13)	12–16(13)	1–3(2)	1
2	4–7(5)	1,2(1)	1,2(1)	1–3(1)	4–7(5)	8–13(11)	13–24(20)
3	1	2–5(3)	1–3(1)	1,2(1)	2–5(3)	6–10(9)	7–14(11)
4	3–7(4)	3,4(3)	2–6(4)	1,2(1)	1,2(1)	2,3(3)	–
5	4–7(5)	3–5(4)	5–9(6)	5–10(8)	7–13(9)	5–8(6)	–
6	18–28(22)	3–5(5)	3–5(4)	3–5(5)	4–7(4)	–	–
7	3–7(5)	4–7(5)	3–6(5)	2–6(4)	3–7(4)	1-S,	5–10(6)
8	1–4(3)	2,3(2)	2–4(3)	2–4(3)	3–6(6)	2-S,	4–12(8)
9	5–9(7)	4–6(5)	4–6(5)	4–7(6)	3–9(6)	6-S,	2–4(3)
10	2–4(3)	2–4(3)	1–3(2)	2–5(4)	4–10(6)	7-S,	1–3(2)
11	2–5(3)	2–5(3)	2,3(2)	2–4(3)	2–4(3)	8-S,	4–7(5)
12	2–4(2)	3–5(3)	2–4(3)	2,3(2)	2–4(2)	9-S,	5–9(7)
13	4–7(5)	4–6(5)	3–6(5)	5–11(8)	2–6(4)	–	–
14	1	1	1	1	1	1	–
15	–	–	–	–	–	–	–

\* Shown in Fig. 2 with fewer than actual branches; nc = not counted.



**FIGURE 2.** Fourth-instar larva of *Anopheles (Cellia) rampae*. A, Head, dorsal (left) and ventral (right) aspects of left side. B, Thorax and abdominal segments I–VI, dorsal (left) and ventral (right) aspects of left side. C, Abdominal segments VII, VIII and X, left side. A, antenna; C, cranium; P, prothorax; M, mesothorax; S, spiracular lobe; T, metathorax; I–VIII, X = abdominal segments I–VIII and X; 1–15 = setal numbers for specified areas, e.g. seta 5-C. Scales in mm.

Crossing experiments involving *An. rampae* (as *An. maculatus* form K), *An. maculatus* forms B and E, *An. sawadwongporni*, *An. dravidicus*, *An. pseudowillmori* and *An. willmori* (Somboon *et al.*, 2008; Thongwat *et al.*, 2008) revealed that *An. rampae* is genetically distinct with varying degrees of genetic incompatibility with the other species. In all crosses, hybrid males were sterile or partially sterile with abnormal and inactive spermatozoa or atrophied testes and accessory glands. In some crosses, male embryos failed to develop or emerge from the eggs. High mortality of larval and/or pupal stages was also observed. One or both ovaries were normal or atrophied in hybrid females. Backcrosses resulted in more severe incompatibility in both sexes. The polytene chromosomes of the ovarian nurse cells in F<sub>1</sub> hybrid females exhibited approximately 70% to complete asynapsis of all chromosome arms.

**Bionomics.** *Anopheles rampae* has been found in sympatry with *An. maculatus*, *An. pseudowillmori* and *An. sawadwongporni*. The immature stages have been found in small rock and sand pools exposed to sunlight, often with green algae, along the Mekong River and in hilly forested areas, about 100–400 m above sea level. Adult females start biting shortly after sunset (18:00–20:00); they are primarily zoophilic but will sometimes bite humans. Whether or not *An. rampae* plays a role in human malaria transmission is unknown.

**Distribution.** *Anopheles rampae* (as *An. maculatus* form K and species K) has been found in localities in northeastern Thailand (Ubon Ratchathani Province – Baimai *et al.*, 1993; Walton *et al.*, 2007; Somboon *et al.*, 2008; Thongwat *et al.*, 2008; Morgan *et al.*, 2009), northern Cambodia (Ratanakiri and Vihear Provinces – Walton *et al.*, 2007; Morgan *et al.*, 2009) and central Vietnam (Quang Binh Province – Walton *et al.*, 2007; Morgan *et al.*, 2009). *Anopheles rampae* has also been found in other areas of northeastern Thailand (Chaiyaphum, Loei, Mukdahan, Nakhon Phanom, Nong Bua Lam Phu, Nong Khai and Phetchabun Provinces) and in Laos (Khammouan Province) (R. Rattanarithikul, unpublished observations). Specimens that form part of the **Type series** (see below) were collected in Udon Thani Province, also in northeastern Thailand.

**Type series.** One hundred and fifty-nine specimens (33 ♀, 23 ♂, 47 Le, 52 Pe, 4 L) derived from four cytotyped progeny broods: UDN 7(3), UDN 7(9), UB 500(14) and UB 501(2). *Holotype*, ♀ (UDN 7(9)-12), with Le and Pe on microscope slide, offspring of female collected as follows, THAILAND: *Udon Thani Province*, Nong Bua Lam Phu (17° 13' N, 102° 27' E), 20 Oct 1986, biting human, coll. Udom. *Paratypes*, same data as holotype: 16 ♀ LePe [UDN 7(9)-1, -10, -13 through -16, -18, -20, -21; UDN 7(3)-2, -3, -5, -12 through -15]; 3 ♀ Pe [UDN 7(3)-1, -16; UDN 7(9)-19]; 4 ♀ [UDN 7(9)-M [mother], UDN 7(9), UDN 7(9)-17; UDN 7(3)-17]; 13 ♂ LePe [UDN 7(9)-2, -4 through -9; UDN 7(3)-6 through -11; 1 ♂ Pe (UDN 7(3)-100]; 1 ♂ [UDN 7(9)]. *Paratypes*, offspring of female collected as follows, THAILAND: *Ubon Ratchathani Province*, Na Chaluai (14° 25' N, 105° 14' E), 7 Oct 1987, biting human, coll. Udom's team: 8 ♀ [UB 501(2)-28, -29, -32, -34, -35; UB 500(14)-21, -22, -100]; 2 ♀ [UB 501(2)-27; UB 500(14)]; 7 ♂ LePe [UB 501(2)-26, -30, -31, -36 through -39]; 1 Pe [UB 501(2)-27]; 4 L [UB 501(2)-A, -B, -C, -D]. The type series is deposited in the Natural History Museum (BMNH), London.

## Acknowledgements

We are grateful to Prachong Panthusiri (formerly of the Armed Forces Research Institute of Medical Sciences, Bangkok) for preparing the illustrations; to the Royal Golden Jubilee PhD Program, the Thailand Research Fund and the Faculty of Medicine Research Fund, Chiang Mai University, Chiang Mai, Thailand, for supporting the PhD studies of Damrongpan Thongwat; and to Prof. Visut Baimai (Mahidol University, Bangkok) and Dr Andrew Polazsek (Natural History Museum, London) for critically reviewing the manuscript.

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